

NREL-Amoco CRADA Phase 3

Bench Scale Report 1.8

Scale-up of Pretreated Corn Fiber Batch SSCF with LNHST2

Project Title: Amoco-NREL CRADA with corn fiber

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Objective

Two batch SSCF (simultaneous saccharification **and** co-fermentation) runs with a mixture of pretreated corn fiber and corn screenings (20% total solids level) were performed to determine how reproducible the bench scale fermentation is **and** how it compares with large scale **PDU** fermentation runs.

Background

This experiment is designed to support the **Task 3 PDU** activities. Results obtained at the bench scale are to **be** compared to data **obtained** at the **PDU** scale to examine scaleability of lab **data** (from 1.7 L to 9,000 L). If the comparison is favorable, then bench scale can be used to collect critical **data** quickly **and** inexpensively.

Materials and Methods

Inoculum

The **inoculum** for each **SSCF** was generated in the **PDU** seed train. That train consisted of five stages starting with cultivation in two 250-mL Erlenmeyer **flask** containing 50 mL of YEPD and progressing to an 160-L fermentor in the **PDU**. The inoculum for the first SSCF was generated with **medium** containing 5% w/v glucose. The first two stages in the laboratory **were** grown in **YPD** (1% w/v yeast extract, 2% w/v peptone **and** 5% w/v glucose, pH 5.0) at 30°C and 150 rpm. The subsequent three stages performed in the **PDU** were grown in 1% w/v corn steep liquor (CSL) with 5% w/v glucose at pH 5.0. The inoculum for the second SSCF was generated in the same manner **as** the first SSCF except 2% w/v glucose was used in each stage instead of 5% w/v glucose due to slower **inoculum** growth associated with the higher **glucose** concentration (see *PDU Run Report, CRADA Task #3*).

Pretreated corn fiber and corn screenings mixture

Pretreated biomass (a mixture of corn fiber and corn screenings) for the first SSCF was prepared in the APR.

The material for the bench-scale SSCF was collected from the APR while PDU vessel 450A was being filled for the first time during PDU Task 3. The pH of this material was adjusted to 5.0 with sodium hydroxide, deionized water was added to bring the solids level down, and an amount of material needed to yield 20% solids (with the addition of enzymes, inoculum, and CSL taken into consideration) in a 1.2-L final volume fermentation was weighed and placed in a 1.7-L fermentor and autoclaved for one hour at 121°C.

The batch 2, SSCF was set-up at the bench scale using material retrieved directly from the PDU (450B, PDU run 4 in Task 3) fermentor after the enzyme, CSL and water had been added to minimize any differences between biomass processing at the bench scale and the PDU. Batch 2 was performed as a compliment to PDU 450B run 4.

Enzyme and nutrients

Cellulase and glucoamylase enzymes (see Report 1.7) were added at 10 IFPU/g cellulose (based on a cellulose content of 17.3% in the raw feed and a cellulase activity of 70 IFPU/mL) and 2 IU/g cellulose (based on a starch content of 17.3% in the raw feed). The enzyme preparation used for the first batch SSCF was filter-sterilized through 0.2 µm filter before being added to the fermentor. Nisin (Alpin and Barrett Ltd, England) was added to both enzyme preparations at 200 mg/L as an antimicrobial agent (to prevent or suppress contamination) before being used in the second batch SSCF.

Corn steep liquor (1% w/v) was added as a nutrient source to each SSCF. To prepare the solution for the first SSCF, the pH of a 50% w/w solution of CSL was adjusted to 5.0 with sodium hydroxide pellets and autoclaved. As mentioned earlier, the material used in batch 2 already contained CSL.

Sampling and Analysis

Initial and final samples were obtained for a total compositional analysis. Samples were also taken during the course of the experiment, and the liquid fraction was analyzed for cellobiose, succinic acid, lactic acid, glycerol, acetic acid, HMF, furfural, and the monomeric and oligomeric sugars glucose, xylose, galactose, arabinose, and mannose. Colony forming units were monitored on each sample to measure cell growth.

Results and Discussions

First Batch SSCF

The major available sugars for use by LNHST2 in this fermentation are monomeric and oligomeric (including cellobiose) glucose in the liquor and cellulose in the solids, as well as the monomeric xylose found in the liquor. Table 1 summarizes the initial and final glucose and xylose levels in both the liquor and the solid fractions in batch 1 and batch 2 SSCF. Within 18.5 hours of SSCF, a majority of the initial monomeric glucose was consumed (Figure 1). Of the total available glucose, 74.09% was consumed in 113 hours. Of the glucan in the solids, 75.4% was either consumed by LNHST2 or converted into oligomeric glucose. Figure 2 shows that the oligomeric concentration increases throughout the fermentation as the glucan level decreases. This demonstrates that

cellulase is actively **breaking down** the cellulose fraction, but that cellobiose is accumulating. The rate of glucose ~~utilization was~~ **3.523 g/L-h**.

After ~~an~~ initial phase of slow uptake of xylose (–12 hours while a majority of glucose was **being** consumed), the rate of **xylose** utilization increased to 0.912 g/L-h. Interestingly, after the monomeric glucose ~~was~~ consumed, the xylose rate decreased to 0.270 g/L-h. By 84 hours, a majority of the monomeric xylose ~~was~~ consumed. By the **end** of the **SSCF**, **92.8%** of the monomeric xylose had been consumed (62.8 1% of the total available xylan).

Table 1: Initial and Final Glucose and Xylose Concentrations and Conversion

		Batch 1			Batch 2		
		t_0	t_f	conversion	b	t_f	conversion
		(g/L)	(g/L)	(%)	(g/L)	(g/L)	(%)
<i>Equivelant Glucose</i>							
soluble	monomeric	50.13	1.69		51.41	1.98	
	oligomeric	10.55	12.39		6.81	9.87	
	cellobiose	2.82	2.79		5.84	1.97	
insoluble	glucan	31.89	7.85	75.38	29.50	6.40	75.30
Total		95.39	24.72	74.09	93.56	20.22	78.39
<i>Equivelant Xylose</i>							
soluble	monomeric	27.96	2.01	92.8 1	26.98	8.62	68.05
	oligomeric	12.08	12.55		8.08	7.01	
insoluble	xylan	1.21	0.78		1.75	0.53	69.9
Total		41.25	15.34	62.81	36.81	16.16	56.10
<i>Glucose and Xylose</i>							
Total		136.64	40.06	70.68	130.37	36.38	72.09

* The duration (t_f) of each run was 113 h for batch 1 and 167 h for batch 2.

During the **SSCF**, 39.74 g/L of ethanol were produced corresponding to ~~an~~ ethanol metabolic yield of 80.5% of theoretical (based on the consumed xylose **and** glucose) (Table 2). The ethanol process yield ~~was~~ 56.910 of theoretical (**based** on the total sugars available). **About 25.4 g/L** of total glucose and monomeric xylose **remained** unused by the end of the fermentation representing a potential for 12.95 g/L more ethanol through process and/or microorganism optimization.

Table 2: Performance Parameters of SSCF by LNHST2

Parameter	Value	
	Batch 1	Batch 2
Glucose conversion	74.09%	78.39%
Xylose Conversion	62.81%	56.10%
Ethanol Process Yield (% theoretical)	56.9%	63.50%
Ethanol Metabolic Yield (% theoretical)	80.5%	88.07%

Table 3 ~~lists~~ the products and their yields based on the consumed ~~sugars~~ in the fermentation. The major by-product ~~was~~ glycerol at 0.045 g/g consumed sugars and xylitol at 0.026 g/g consumed sugars. The cell ~~mass~~ yield ~~was~~ based on an estimated yield of -0.05 g/g consumed glucose; since cell ~~mass~~ quantification in the presence of insoluble solids is difficult and unreliable. Previous studies on ~~pure~~ sugars indicated that the cell ~~mass~~ yield can ~~vary~~ from 0.04 to 0.10 g/g of consumed sugars. The overall ~~mass~~ balance closure was 92.37%.

Table 3: By-product and Carbon Balance for Batch 1 and 2 SSCF

g product 100 g consumed glucose and xylose		
Products	Batch 1	Batch 2
Ethanol	41.14	45.00
Cell Mass	4.04	3.94
Carbon Dioxide	39.35	43.05
Glycerol	4.47	5.02
Aceuc Acid	0.32	0.00
Lactic Acid	0.00	0.17
Succinic Acid	0.45	0.55
Xylitol	2.61	2.75
Total	92.37	100.48

Second Batch SSCF

A second batch SSCF ~~was~~ performed with material processed in the PDU (pH adjusted, CSL ~~and~~ enzymes added, and water added to bring the total ~~solids~~ to the appropriate operating level) to ~~minimize~~ any differences between the material used ~~for~~ bench scale and PDU experiments. The pretreatment process liberated 64.06 g/L of soluble glucose (monomeric and oligomeric) ~~and~~ **26.98** g/L of monomeric xylose, which are similar to the levels in batch 1 (63.50 g/L ~~and~~ 27.96 g/L, respectively). After a lag of 6 h, glucose ~~was~~ used at a rate of **2.41** g/L-h. Within 24 h of inoculation, all of the monomeric glucose ~~was~~ consumed (Figure 3). After that, the rate-limiting step was the liberation of glucose ~~from~~ oligomers ~~and~~ cellulose by the catalytic action of cellulase. The overall glucose conversion, including the total soluble and insoluble glucan fractions, was 78.39% after **167** hours in batch mode (Table 1).

Of the 26.98 g/L monomeric xylose available, 18.36 g/L was consumed, representing a conversion of 68.1% (Table 1) (**56.7%** conversion of the total xylan). The conversion rate for xylose was 0.249 g/L-h. The phenomena observed in the first batch, where xylose consumption was higher in the presence of glucose, was not apparent in batch 2. It should be noted that in the batch 1 SSCF, 92.8% of the monomeric xylose was consumed. The major difference between the two studies was the amount of acetic acid produced during pretreatment (2.52 g/L in batch 1 compared to 4.93 g/L in batch 2) (Figure 4). The inhibitory effect of high acetic acid concentration on xylose utilization and ethanol production has been confirmed by studies performed with a variety of yeast strains at various pH levels (Van Zyl et al 1991)*. Hence, the incomplete use of monomeric xylose by LNHST2, as evidenced by the process yield of 63.50%, could be attributed to the detrimental effect of acetic acid inhibition (and perhaps other inhibitors) on cell metabolism. Nevertheless, even though not all the xylose was consumed, the ethanol metabolic yield was high, 88.07% (Table 2) or 0.450 g ethanol per gram of consumed sugars.

In addition to the decrease in xylose utilization, the cell concentration was also lower in Batch 2 compared to Batch 1 (Figure 5). The cell mass in batch 1 increased to 1.07×10^8 cells/mL by 18.5 h, where it remained during the fermentation. In batch 2, the cell population reached a maximum of 6.8×10^7 cells/mL at 25 h into the fermentation and then declined. Again, inhibition of cell metabolism may be the reason.

The distribution of by-products was very similar to that observed in batch 1. As shown in Table 3, glycerol was the major by-product at 0.050 g/g consumed sugars. The overall mass balance closure was very satisfactory at 100.48%.

Correlation between PDU and bench scale run

The pretreated biomass was also subjected to SSCF in a 9,000-L pilot plant fermentor under conditions similar to those at the bench scale. As seen in Figure 3, the results of the pilot plant run correlated very well with the bench-scale data. After 24 hours of SSCF, the concentrations of glucose, xylose, and ethanol were 2.7 g/L, 24.9 g/L, and 29.4 g/L in the bench scale fermentor and 3.7 g/L, 23.8 g/L, and 31.1 g/L in the pilot fermentor. At 96 hours, the concentrations of these same components were 1.5 g/L, 11.9 g/L, and 40.5 g/L in the bench reactor and 1.7 g/L, 9.4 g/L, and 43.4 g/L in the pilot-scale reactor. Similarly, by-product formation was in close agreement, as indicated by xylitol, which reached 2.4-2.5 g/L in both vessels (data not shown). The good correlation shows that despite the 5,300-fold increase in fermentor size, the performance of ST2 remained unchanged and very promising. The scalability and reproducibility of the SSCF performance justify further optimization studies for use of the recombinant cofermenting yeast in commercial applications.

Conclusions

The data obtained from the two bench scale fermentations correlate quite well, as the total glucose and xylose consumption was 71.87% and 72.98% in batch 1 and batch 2, respectively. The slightly lower xylose consumption in batch 2 seems to be the result of cell inhibition by acetic acid and ethanol. The presence of other inhibitors and possible nutrient limitations may have also played a role in that phenomenon. Finally, the data from the bench scale batch 2 and the PDU 450B run 4 correlate very well indicating that the bench scale results are readily scaleable.

*Van Zyl, C., Prior, B. A., Du Preez, J., *Enzyme Microb. Technol.* 13, 82-86. (1991).

Figure 1: SSCF Batch 1 with LNHST2
(Task 3 Bench Scale Data Run 1, PDU Run 1)

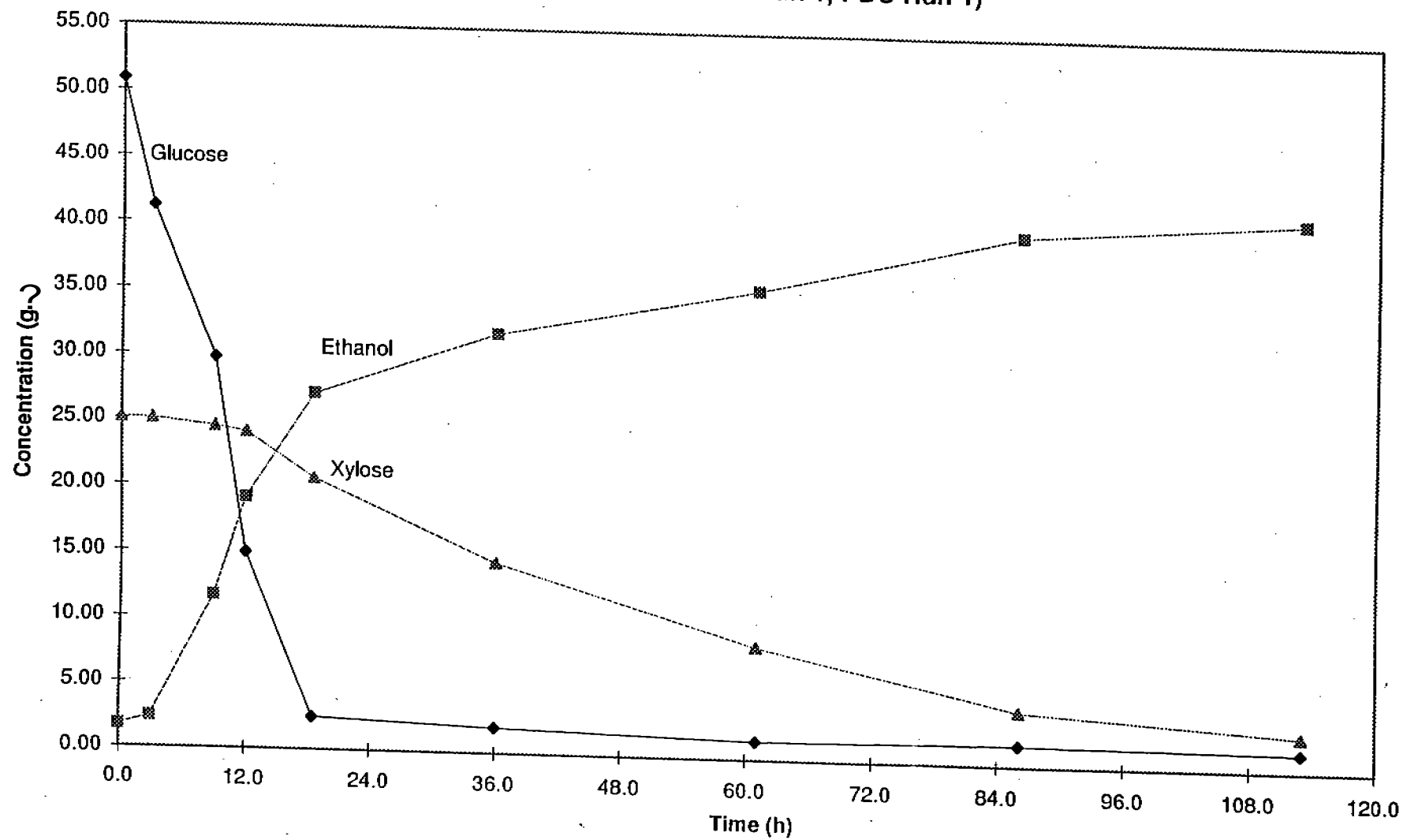


Figure 2: Batch 1 SSCF Sugar Consumption

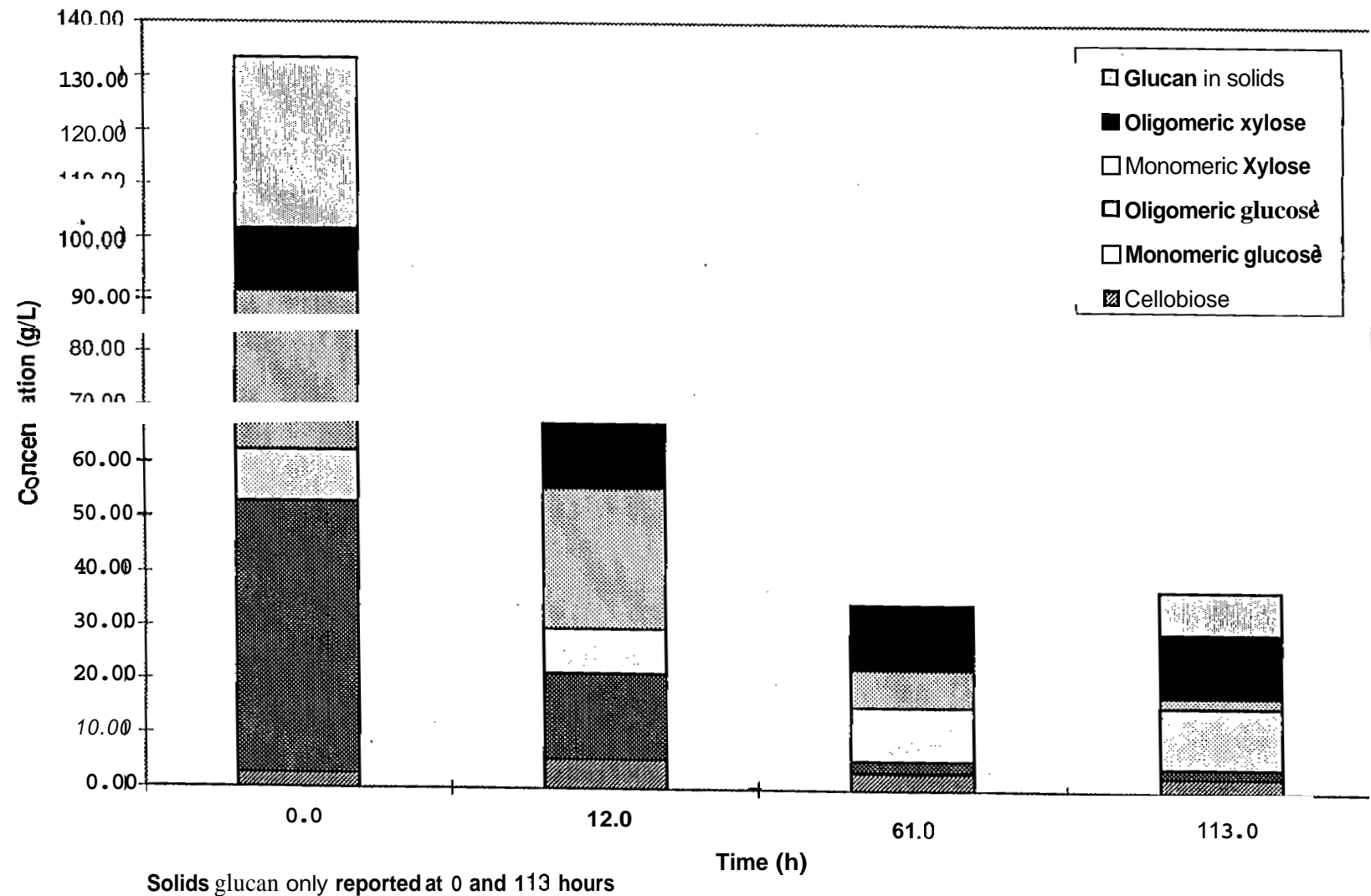
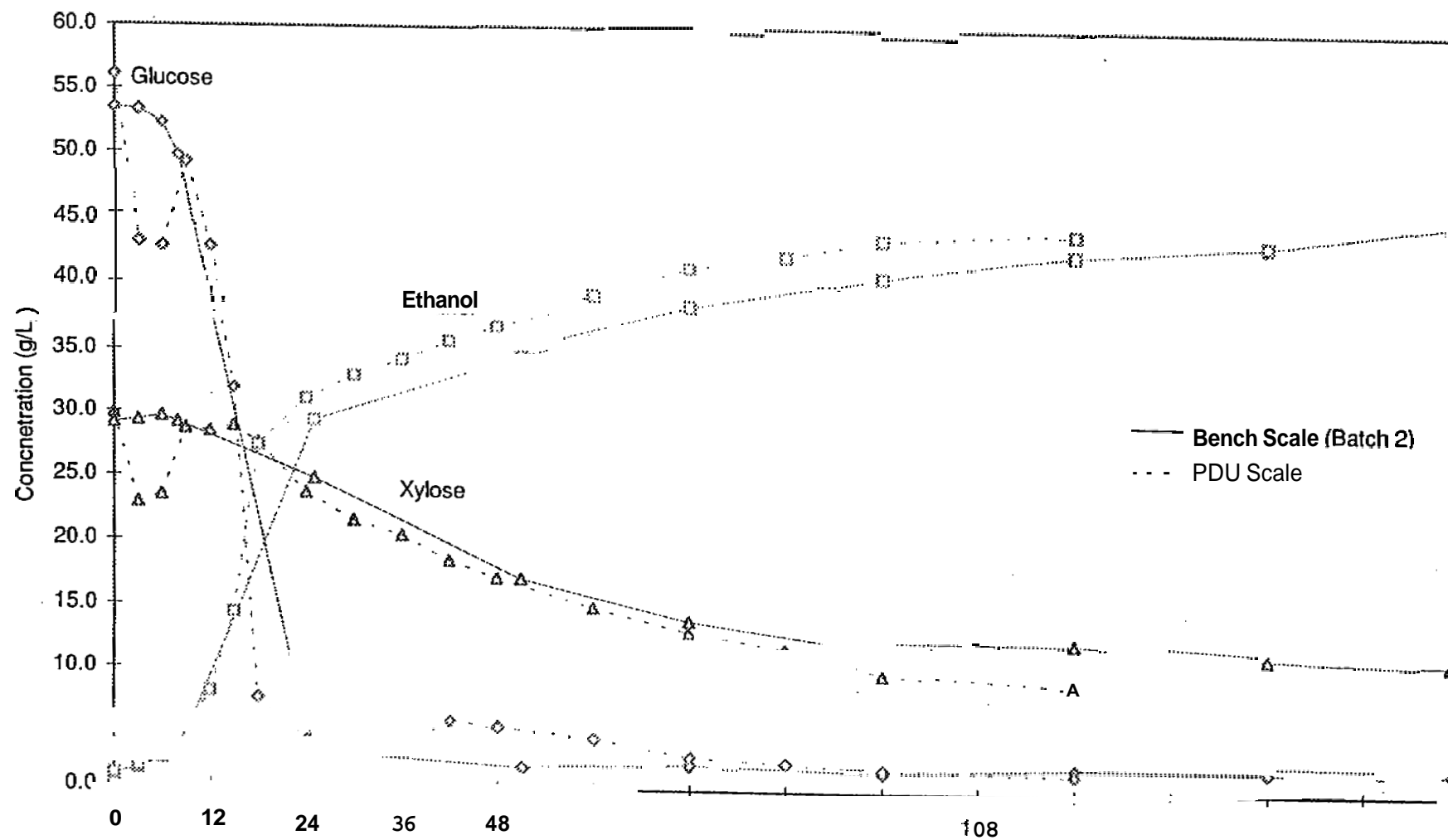


Figure3: Comparison of Bench Scale and PDU Scale SSCF Results with LNHST2



**Figure 4: By-products from SSCFs with LNHST2
(Comparison of Bench Scale Data Runs 1 and 2)**

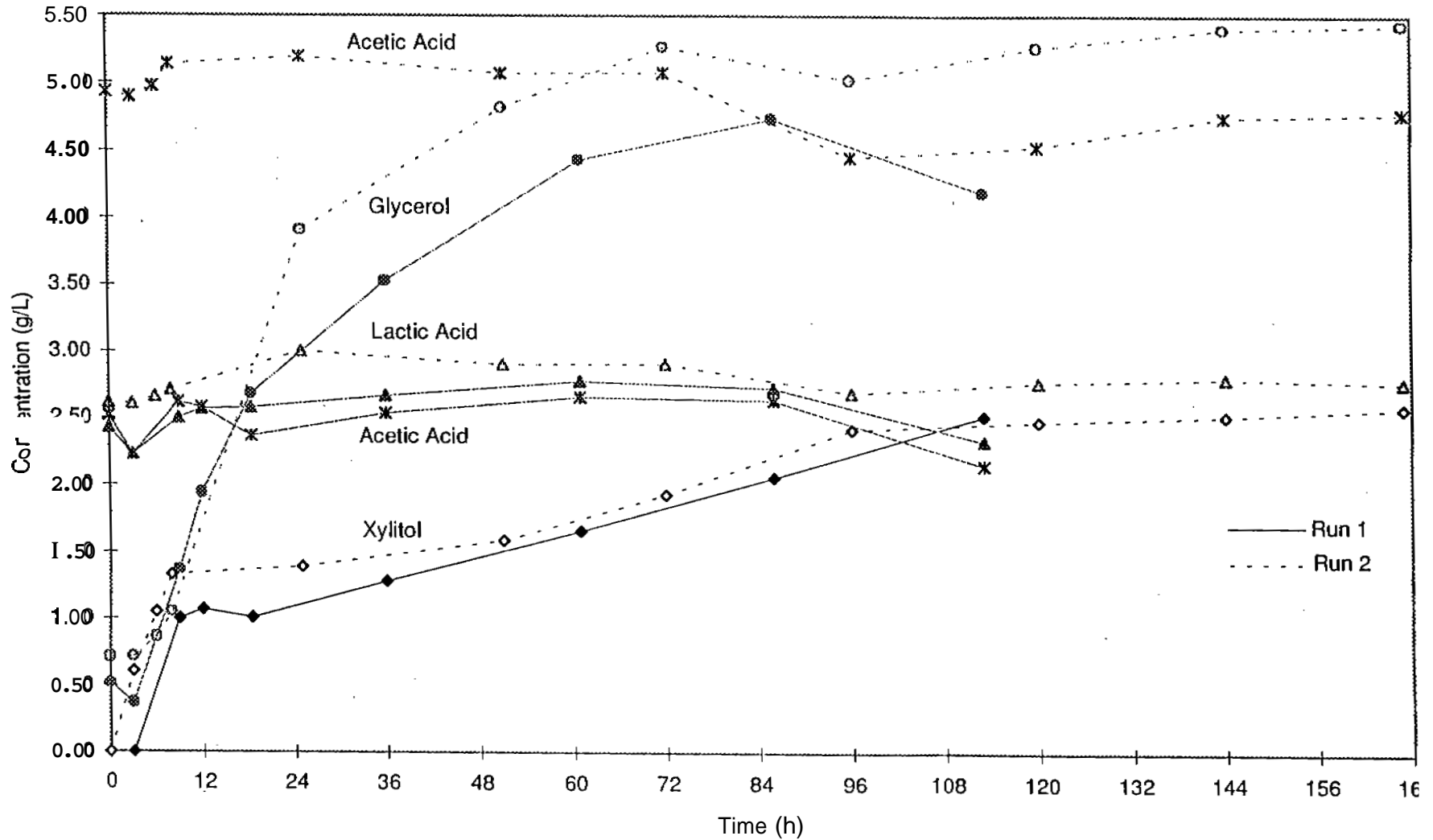
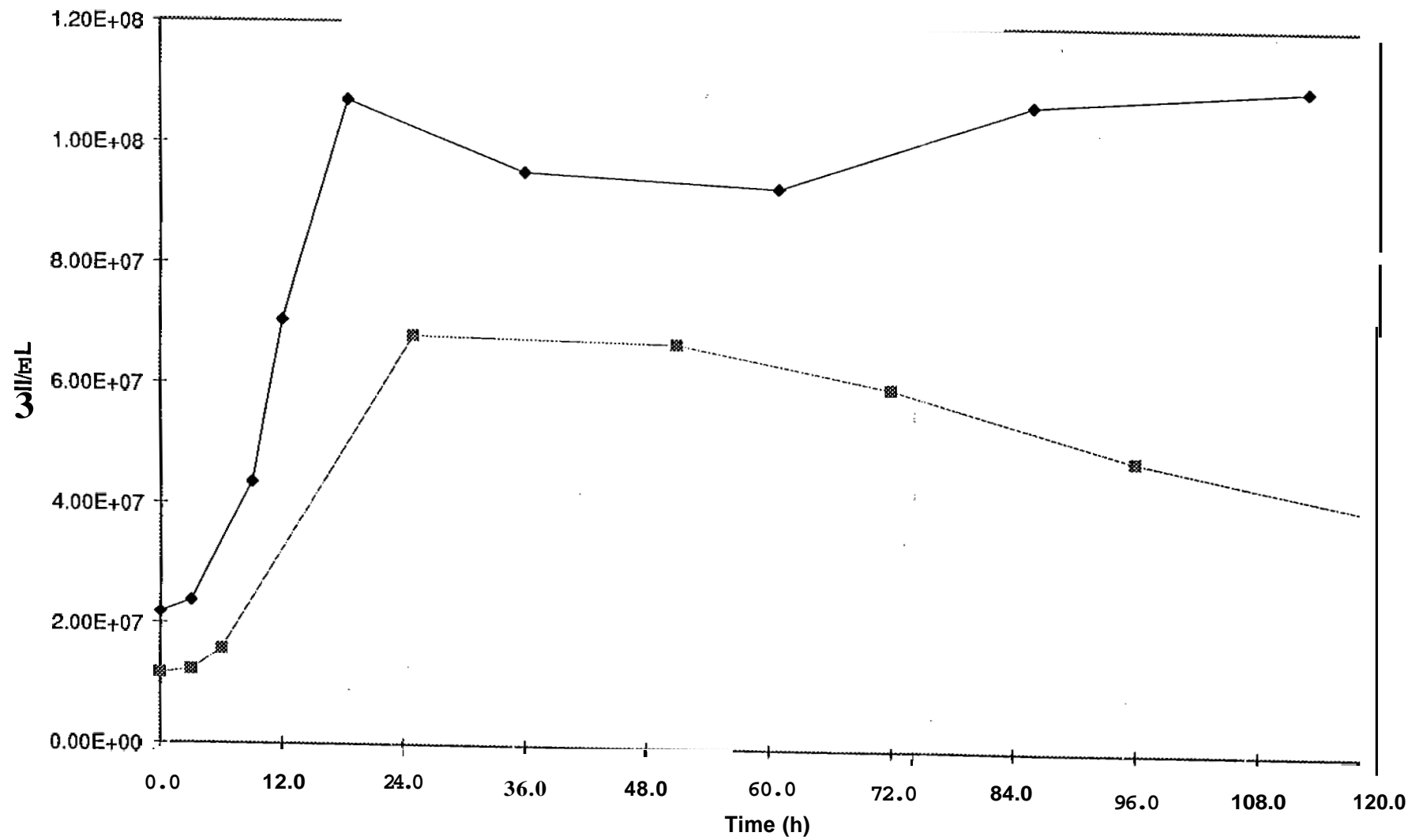


Figure 5: Comparison of Colony Forming Units Between Batch One and Two SSCF



Raw Data

(Batch 2 in Report 1.0) CAT Task Analysis No. 96-028 (liquors) and 96-029 (solids)

Tracking samples performed by NREL

By-products

Sample	Time (h)	Glucose	Glucose	Xylose	Ethanol	Ethanol	Ethanol	Cellobiose	Xylitol	Succinic acid	Lactic acid	Glycerol	Acetic acid	Ethanol	CFU/mL
		(g/L)	(g/L)	(g/L)	(g/L)	(g/L)	(g/L)	(g/L)							
1	0	48.88	53.45	29.12	1.36	0.92	1.10	5.36	0.00	0.00	2.61	0.71	4.93	1.10	1.17E+07
2	3	50.00	53.31	29.33	1.88	1.20	1.40	5.74	0.61	0.19	2.61	0.72	4.89	1.40	1.22E+07
3	6	47.88	52.25	29.63	2.58	2.09	2.10	6.08	1.05	0.28	2.66	0.87	4.97	2.10	1.57E+07
4	8	46.63	49.70	29.16	3.13	2.89	2.90	6.21	1.33	0.56	2.71	1.05	5.13	2.90	
5	25	0.60	2.70	24.86	28.63	29.39	27.40	3.99	1.39	0.61	3.00	3.91	5.19	27.40	6.80E+07
6	51	0.46	1.71	17.21	33.13	34.66	36.60	3.02	1.59	0.55	2.90	4.82	5.07	36.60	6.70E+07
7	72	0.45	1.96	14.03	37.21	30.07	36.00	2.65	1.93	0.71	2.91	5.26	5.07	36.00	6.00E+07
8	96	0.45	1.53	11.85	38.63	40.48	30.60	2.30	2.42	0.47	2.69	5.02	4.45	38.60	4.83E+07
9	120	0.41	2.00	11.73	39.50	42.40	40.20	2.47	2.48	0.51	2.77	5.26	4.53	40.20	4.00E+07
10	144	0.41	1.99	10.79	39.90	43.25	41.50	2.44	2.52	0.49	2.80	5.40	4.75	41.50	1.70E+07
11	167	0.39	1.88	10.45	39.82	45.09	43.40	2.34	2.58	0.52	2.77	5.43	4.78	43.40	1.40E+07

(Batch 1 In Report 1.8) CAT Task Analysis No. 96-025 (liquors) and 96-027 (solids)

Tracking samples performed by NREL

By-products

Sample	Time (h)	Glucose	Glucose	Xylose	Ethanol	Ethanol	Ethanol	Cellobiose	Xylitol	Succinic acid	Lactic acid	Glycerol	Acetic acid	Ethanol	CFU/mL
		(g/L)	(g/L)	(g/L)	(g/L)	(g/L)	(g/L)	(g/L)							
1	0.0	42.50	50.87	25.10	1.90	1.69	0.90	2.68	0.00	0.00	2.43	0.52	2.52	0.90	2.18E+07
2	3.0	43.90	41.21	25.05	2.77	2.34	2.60	3.00	0.00	0.00	2.23	0.37	2.23	2.60	2.37E+07
3	9.0	31.00	29.75	24.53	13.10	11.68	11.40	4.05	1.00	0.31	2.50	1.37	2.62	11.40	4.35E+07
4	12.0	11.40	14.97	24.09	20.80	19.12	18.30	4.26	1.07	0.36	2.57	1.94	2.58	18.30	7.05E+07
5	18.5	0.40	2.46	20.67	27.80	27.10	25.60	3.11	1.01	0.34	2.58	2.68	2.37	25.60	1.07E+08
6	36.0	0.37	1.94	14.50	31.58	31.86	29.80	2.46	1.28	0.42	2.67	3.53	2.54	29.80	9.53E+07
7	61.0	0.23	1.41	8.55	38.34	35.54	33.30	2.63	1.66	0.46	2.78	4.43	2.66	33.30	9.30E+07
8	86.0	0.20	1.61	4.13	39.98	40.01	36.90	1.50	2.06	0.45	2.73	4.74	2.64	36.90	1.07E+08
9	113.0	0.18	1.39	2.67	39.22	41.42	37.90	1.52	2.52	0.42	2.33	4.19	2.15	37.90	1.10E+08

Raw Data

Batch 2

Monomeric and Oligomeric Sugars Analysis												
Cellobiose	Monomeric Glucose	Oligomeric Glucose	Total soluble glucose	Monomeric Xylose	Oligomeric Xylose	Total soluble xylose	Monomeric galactose	Oligomeric galactose	Total soluble galactose	Monomeric arabinose	Oligomeric arabinose	Total soluble arabinose
(g/L)	(g/L)	(g/L)	(g/L)	(g/L)	(g/L)	(g/L)	(g/L)	(g/L)	(g/L)	(g/L)	(g/L)	(g/L)
5.55	51.41	6.13	63.09	26.98	7.57	34.55	6.54	0.59	7.13	20.75	1.08	21.83
5.96	50.44			26.95			6.79			20.60		
6.21	48.97			26.46			6.48			20.54		
6.39	47.46			26.78			6.53			20.74		
3.79	4.52	10.48	18.79	22.71	7.98	30.69	6.56	0.98	7.54	18.72	2.32	21.04
2.53	3.10	11.00	16.63	16.00	7.81	23.89	6.82	0.66	7.48	18.18	2.31	20.49
2.43	2.73	11.58	16.74	12.67	7.31	19.98	7.09	-0.03	7.06	18.17	1.76	19.93
2.08	2.51	10.44	15.03	10.50	7.62	18.12	4.67	1.12	5.79	17.46	3.61	21.07
2.03	2.29	9.92	14.24	9.65	7.21	16.86	4.75	1.00	5.75	17.42	3.43	20.85
1.94	0.44	11.25	13.63	8.97	7.12	16.09	4.82	0.85	5.67	17.20	4.36	21.56
1.87	1.98	8.88	12.73	8.62	6.56	15.18	4.85	0.68	5.53	17.07	3.65	20.72

Batch 1

Monomeric and Oligomeric Sugars Analysis												
Cellobiose	Monomeric Glucose	Oligomeric Glucose	Total soluble glucose	Monomeric Xylose	Oligomeric Xylose	Total soluble xylose	Monomeric galactose	Oligomeric galactose	Total soluble galactose	Monomeric arabinose	Oligomeric arabinose	Total soluble arabinose
(g/L)	(g/L)	(g/L)	(g/L)	(g/L)	(g/L)	(g/L)	(g/L)	(g/L)	(g/L)	(g/L)	(g/L)	(g/L)
2.68	50.13	9.50	62.31	27.96	11.31	39.27	5.93			18.21	5.04	23.25
5.93	40.2			27.53			5.95			17.87		
5.98	29.72			26.86			5.83			17.50		
5.4t	15.78	8.36	29.55	26.01	11.86	37.87	5.75			16.99	3.43	20.42
3.58	4.92			18.61			5.45			15.27		
3.16	3.73	7.48	14.37	12.72	6.11	18.83	5.15	0.79	5.94	15.08	1.38	16.46
3.19	2.2	9.83	15.22	7.01	12.15	19.16	5.10	0.52	5.62	14.33	4.15	18.48
3.06	2.03			3.04			5.09			14.32		
2.65	1.69	11.15	15.49	2.01	11.75	13.76	4.95	0.17	5.12	13.96	3.86	17.82

Raw Data

Batch 2

Monomeric mannose (g/L)	Total soluble mannose (g/L)
0.00	0.05
0.00	
0.00	
0.00	
0.00	0.11
0.00	0.08
0.00	0.09
0.00	0.07
0.00	0.06
0.00	0.09
0.00	0.10

Batch 1

Monomeric mannose (g/L)	Total soluble mannose (g/L)
0.00	0.00
0.00	
0.00	
0.00	0.00
0.00	
0.00	0.00
0.00	0.00
0.00	
0.00	0.00

Solids Analysis (% Dry weight)

Glucose	Xylose	Galactose	Arabinose	Mannose	Klason Lignin	Acid Soluble Lignin	Total Ash	Total Solids	HMF (g/L)	Furfural (g/L)
40.18	2.38	0.40	2.04	0.00	32.99	7.46	1.67	22.20	0.27	0.33
									0.26	0.28
									0.25	0.19
									0.22	0.14
									0.00	0.14
									0.00	0.73
									0.00	0.76
									0.00	0.00
									0.00	0.00
13.40	1.10	0.20	0.80	0.00	46.81	10.29	0.52	22.46	0.00	0.00

Solids Analysis (% Dry weight)

Glucose	Xylose	Galactose	Arabinose	Mannose	Klason Lignin	Acid Soluble Lignin	Total Ash	Total Solids	HMF (g/L)	Furfural (g/L)
55.2	2.1	0.1	2.0	0.0	20.1	6.69	0.3	11.2	0.22	0.18
									0.21	0.13
									0.00	0.00
									0.09	0.00
									0.00	0.00
									0.08	0.00
									0.00	0.00
									0.00	0.00
20.62	2.04	0.15	1.04	0.00	3636.00	8.39	0.90	22.01	0.00	0.00